

SEQUENCE LISTING

<110> Le, Junming Vilcek, Jan Daddona, Peter Ghrayeb, John Knight, David M. Siegel, Scott

<120> Anti-TNF Antibodies and Peptides of Human Tumor Necrosis Factor

<130> 0975.1005-006

<140> US 09/756,398

<141> 2001-01-08

<150> U.S. 09/133,119

<151> 1998-08-12

<150> U.S. 08/570,674

<151> 1995-12-11

<150> U.S. 08/324,799

<151> 1994-10-18

<150> U.S. 08/192,102

<151> 1994-02-04

<150> U.S. 08/192,861

<151> 1994-02-04

<150> U.S. 08/192,093

<151> 1994-02-04

<150> U.S. 08/010,406

<151> 1993-01-29

<150> U.S. 08/013,413

<151> 1993-02-02

<150> U.S. 07/943,852

<151> 1992-09-11

<150> U.S. 07/853,606

<151> 1992-03-18

<150> U.S. 07/670,827

<151> 1991-03-18

<160> 19

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 157

<212> PRT

<213> Homo sapiens

<400> 1 Val Arg Ser Ser Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val 10 Val Ala Asn Pro Gln Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg 25 Ala Asn Ala Leu Leu Ala Asn Gly Val Glu Leu Arg Asp Asn Gln Leu Val Val Pro Ser Glu Gly Leu Tyr Leu Ile Tyr Ser Gln Val Leu Phe 55 Lys Gly Gln Gly Cys Pro Ser Thr His Val Leu Leu Thr His Thr Ile 70 75 Ser Arg Ile Ala Val Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala 90 Ile Lys Ser Pro Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys 100 105 Pro Trp Tyr Glu Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys 115 120 125 Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg Pro Asp Tyr Leu Asp Phe 130 Ala Glu Ser Gly Gln Val Tyr Phe Gly Ile Ile Ala Leu 150

<210> 2 <211> 321 <212> DNA <213> Mus Balb/c <220> <221> CDS

<221> CDS <222> (1)...(321)

gac atc ttg ctg act cag tct cca gcc atc ctg tct gtg agt cca gga 48
Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
1 5 10 15

gaa aga gtc agt ttc tcc tgc agg gcc agt cag ttc gtt ggc tca agc 96 Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser 20 25 30

atc cac tgg tat cag caa aga aca aat ggt tct cca agg ctt ctc ata 144
Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
35 40 45

aag tat gct tct gag tct atg tct ggg atc cct tcc agg ttt agt ggc 192 Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly 50 55 60

agt gga tca ggg aca gat ttt act ctt agc atc aac act gtg gag tct 240 Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser 65 70 75 80

gaa gat att gca gat tat tac tgt caa caa agt cat agc tgg cca ttc 288
Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
85

321

acg ttc ggc tcg ggg aca aat ttg gaa gta aaa

Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys

```
100
<210> 3
<211> 107
<212> PRT
<213> Mus Balb/c
<400> 3
Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser
            20
                               25
                                                   30
Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
        35
                           40
Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly
   50
                       55
                                           60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser
               Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
               85
                                   90
Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys
            100
                               105
<210> 4
<211> 357
<212> DNA
<213> Mus Balb/c
<220>
<221> CDS
<222> (1)...(357)
<400> 4
gaa gtg aag ctt gag gag tct gga gga ggc ttg gtg caa cct gga gga
Glu Val Lys Leu Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
                5
                                                        15
tcc atg aaa ctc tcc tgt gtt gcc tct gga ttc att ttc agt aac cac
Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His
            20
tgg atg aac tgg gtc cgc cag tct cca gag aag ggg ctt gag tgg gtt
Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val
        35
gct gaa att aga tca aaa tct att aat tct gca aca cat tat gcq qaq
Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu
    50
                        55
tct gtg aaa ggg agg ttc acc atc tca aga gat gat tcc aaa agt gct
Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala
65
                    70
                                        75
gtc tac ctg caa atg acc gac tta aga act gaa gac act ggc gtt tat
```

Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr

357

85 90 95

tac tgt tcc agg aat tac tac ggt agt acc tac gac tac tgg ggc caa 336
Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln
100 105 110

ggc acc act ctc aca gtc tcc Gly Thr Thr Leu Thr Val Ser 115

<210> 5 <211> 119 <212> PRT <213> Mus Balb/c

<400> 5

Glu Val Lys Leu Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His

Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val 35 40 45

Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu 50 55 60

Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala 70 75 80

Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr 85 90 95

Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln 100 105 110

Gly Thr Thr Leu Thr Val Ser 115

<210> 6

<211> 8

<212> PRT

<213> Homo sapiens

<400> 6

Gly Thr Leu Val Thr Val Ser Ser

<210> 7

<211> 7

<212> PRT

<213> Homo sapiens

<400> 7

Gly Thr Lys Leu Glu Ile Lys
1 5

<210> 8

<211> 20

<212> DNA

<213> Artificial Sequence

<220> <223> PCR oligonucleotides	
<400> 8 cctggatacc tgtgaaaaga	20
<210> 9 <211> 27 <212> DNA	
<213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 9 cctggtacct tagtcaccgt ctcctca	27
<210> 10 <211> 27 <212> DNA	. a mos vol. a somens of p
<213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 10 aatagatatc tccttcaaca cctgcaa	27
<210> 11 <211> 21 <212> DNA <213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 11 atcgggacaa agttggaaat a	21
<210> 12 <211> 16 <212> DNA <213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 12 ggcggtctgg taccgg	16
<210> 13 <211> 19 <212> DNA <213> Artificial Sequence	

<220> <223> PCR oligonucleotides	
<400> 13 gtcaacaaca tagtcatca	19
<210> 14 <211> 23 <212> DNA <213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 14 cacaggtyty tececaagga aaa	23
<210> 15 <211> 18 <212> DNA <213> Artificial Sequence	ren emana eman
<220> <223> PCR oligonucleotides	
<400> 15 aatctggggt aggcacaa	18
<210> 16 <211> 17 <212> DNA <213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 16 agtgtgtgtc cccaagg	17
<210> 17 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> PCR oligonucleotides	
<400> 17 cacagetgee egeceaggtg geat	24
<210> 18	

4

<211><212><213>	
<220> <223>	PCR oligonucleotides
<400> gtcgc	18 Cagtg ctccctt
<210><211><211><212><213>	20
	PCR oligonucleotides

<400> 19 atcggacgtg gacgtgcaga

20

17

ŧ